This Page Is Inserted by IFW Operations
and is not a part of the Official Record

BEST AVAILABLE IMAGES

Defective images within this document are accurate representations of
the original documents submitted by the applicant.

Defects in the images may include (but are not limited to):

- BLACK BORDERS

- TEXT CUT OFF AT TOP, BOTTOM OR SIDES

- FADED TEXT

- ILLEGIBLE TEXT

- SKEWED/SLANTED IMAGES

- COLORED PHOTOS

- BLACK OR VERY BLACK AND WHITE DARK PHOTOS

- GRAY SCALE DOCUMENTS

IMAGES ARE BEST AVAILABLE COPY.

As rescanning documents *will not* correct images,
please do not report the images to the
Image Problem Mailbox.

M.I.C

0400
4-25-01
#2

OIPE

ENTERED

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/820,788

DATE: 04/10/2001
TIME: 15:48:01

Input Set : A:\Seqlist.txt
Output Set: N:\CRF3\04102001\I820788.raw

```
 4 <110> APPLICANT: SHAO, Wei et al
 6 <120> TITLE OF INVENTION: ISOLATED HUMAN DRUG-METABOLIZING
 7       PROTEINS, NUCLEIC ACID MOLECULES ENCODING HUMAN
 8       DRUG-METABOLIZING PROTEINS,
 9       AND USES THEREOF
12 <130> FILE REFERENCE: CL001207
14 <140> CURRENT APPLICATION NUMBER: US/09/820,788
14 <141> CURRENT FILING DATE: 2001-03-30
14 <160> NUMBER OF SEQ ID NOS: 4
16 <170> SOFTWARE: FastSEQ for Windows Version 4.0
18 <210> SEQ ID NO: 1
19 <211> LENGTH: 1537
20 <212> TYPE: DNA
21 <213> ORGANISM: Human
23 <400> SEQUENCE: 1
24 cctgcctggt cctctgtgcc tggtgggggtg ggggtgccag gtgtgtccag aggagcccat   60
25 ttggtagtga ggcaggtatg gggctagaag cactggtgcc cctggccgtg atagtggcca  120
26 tcttcctgct cctggtggac ctgatgcacc ggcgccaacg ctgggctgca cgctactcac  180
27 caggccccct gccactgccc gggctgggca acctgctgca tgtggacttc cagaacacac  240
28 catactgctt cgaccagttg cggcgccgct tcgggacgt gttcagcctg cagctggcct  300
29 ggacgccggt ggtcgtgctc aatggctggg cggccgtgcg cgaggcgctg gtgacccacg  360
30 gcgaggacac cgccgaccgc ccgcctgtgc ccatcaccca gatcctgggt tttgggccgc  420
31 gttcccaagg acgccccttt cgccccaacg gtctcttgga caaagccgtg agcaacgtga  480
32 tcgcctccct cacctgcggg cgccgcttcg agtacgacga ccctcgcttc ctcaggctgc  540
33 tggacctagc tcaggaggga ctgaaggagg agtcgggctt tctgcgcgag gtgctgaatg  600
34 ctgtccccgt cctcctgcat atcccagcgc tggctggcaa ggtcctacgc ttccaaaagg  660
35 ctttcctgac ccagctggat gagctgctaa ctgagcacag gatgacctgg gacccagccc  720
36 agccccccccg agacctgact gaggccttcc tggcagagat ggagaaggcc aaggggaacc  780
37 ctgagagcag cttcaatgat gagaacctgc gcatagtggt ggctgacctg ttctctgccg  840
38 ggatggtgac cacctcgacc acgctggcct ggggcctcct gctcatgatc ctacatccgg  900
39 atgtgcagcg ccgtgtccaa caggagatcg acgacgtgat agggcaggtg cggcgaccag  960
40 agatgggtga ccaggctcac atgccctaca ccactgccgt gattcatgag gtgcagcgct 1020
41 ttggggacat cgtcccccctg ggtgtgaccc atatgacatc ccgtgacatc gaagtacagg 1080
42 gcttccgcat ccctaaggga acgacactca tcaccaacct gtcatcggtg ctgaaggatg 1140
43 aggccgtctg ggagaagccc ttccgcttcc accccgaaca cttcctggat gcccagggcc 1200
44 actttgtgaa gccggaggcc ttcctgcctt tctcagcagg ccgccgtgca tgcctcgggg 1260
45 agcccctggc ccgcatggag ctcttcctct tcttcacctc cctgctgcag cacttcagct 1320
46 tctcggtgcc cactggacag ccccggccca gccaccatgg tgtcttgct ttcctggtga 1380
47 ccccatcccc ctatgagctt tgtgctgtgc cccgctagaa tggggtacct agtccccagc 1440
48 ctgctcccta gccagaggct ctaatgtaca ataaagcaat gtggtagttc caaaaaaaaa 1500
49 aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaa                          1537
51 <210> SEQ ID NO: 2
52 <211> LENGTH: 446
53 <212> TYPE: PRT
54 <213> ORGANISM: Human
56 <400> SEQUENCE: 2
57 Met Gly Leu Glu Ala Leu Val Pro Leu Ala Val Ile Val Ala Ile Phe
```

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/820,788

DATE: 04/10/2001
TIME: 15:48:01

Input Set : A:\Seqlist.txt
Output Set: N:\CRF3\04102001\I820788.raw

```
58  1              5                   10                  15
59  Leu Leu Leu Val Asp Leu Met His Arg Arg Gln Arg Trp Ala Ala Arg
60              20                  25                  30
61  Tyr Ser Pro Gly Pro Leu Pro Leu Pro Gly Leu Gly Asn Leu Leu His
62          35                  40                  45
63  Val Asp Phe Gln Asn Thr Pro Tyr Cys Phe Asp Gln Leu Arg Arg Arg
64          50                  55                  60
65  Phe Gly Asp Val Phe Ser Leu Gln Leu Ala Trp Thr Pro Val Val Val
66  65                  70                  75                  80
67  Leu Asn Gly Leu Ala Ala Val Arg Glu Ala Leu Val Thr His Gly Glu
68              85                  90                  95
69  Asp Thr Ala Asp Arg Pro Pro Val Pro Ile Thr Gln Ile Leu Gly Phe
70              100                 105                 110
71  Gly Pro Arg Ser Gln Gly Arg Pro Phe Arg Pro Asn Gly Leu Leu Asp
72          115                 120                 125
73  Lys Ala Val Ser Asn Val Ile Ala Ser Leu Thr Cys Gly Arg Arg Phe
74          130                 135                 140
75  Glu Tyr Asp Asp Pro Arg Phe Leu Arg Leu Leu Asp Leu Ala Gln Glu
76  145                 150                 155                 160
77  Gly Leu Lys Glu Glu Ser Gly Phe Leu Arg Glu Val Leu Asn Ala Val
78              165                 170                 175
79  Pro Val Leu Leu His Ile Pro Ala Leu Ala Gly Lys Val Leu Arg Phe
80              180                 185                 190
81  Gln Lys Ala Phe Leu Thr Gln Leu Asp Glu Leu Leu Thr Glu His Arg
82          195                 200                 205
83  Met Thr Trp Asp Pro Ala Gln Pro Pro Arg Asp Leu Thr Glu Ala Phe
84      210                 215                 220
85  Leu Ala Glu Met Glu Lys Ala Lys Gly Asn Pro Glu Ser Ser Phe Asn
86  225                 230                 235                 240
87  Asp Glu Asn Leu Arg Ile Val Val Ala Asp Leu Phe Ser Ala Gly Met
88              245                 250                 255
89  Val Thr Thr Ser Thr Thr Leu Ala Trp Gly Leu Leu Leu Met Ile Leu
90              260                 265                 270
91  His Pro Asp Val Gln Arg Arg Val Gln Gln Glu Ile Asp Asp Val Ile
92          275                 280                 285
93  Gly Gln Val Arg Arg Pro Glu Met Gly Asp Gln Ala His Met Pro Tyr
94      290                 295                 300
95  Thr Thr Ala Val Ile His Glu Val Gln Arg Phe Gly Asp Ile Val Pro
96  305                 310                 315                 320
97  Leu Gly Val Thr His Met Thr Ser Arg Asp Ile Glu Val Gln Gly Phe
98              325                 330                 335
99  Arg Ile Pro Lys Gly Thr Thr Leu Ile Thr Asn Leu Ser Ser Val Leu
100             340                 345                 350
101 Lys Asp Glu Ala Val Trp Glu Lys Pro Phe Arg Phe His Pro Glu His
102         355                 360                 365
103 Phe Leu Asp Ala Gln Gly His Phe Val Lys Pro Glu Ala Phe Leu Pro
104     370                 375                 380
105 Phe Ser Ala Gly Arg Arg Ala Cys Leu Gly Glu Pro Leu Ala Arg Met
106 385                 390                 395                 400
```

```
107 Glu Leu Phe Leu Phe Phe Thr Ser Leu Leu Gln His Phe Ser Phe Ser
108                 405                 410                 415
109 Val Pro Thr Gly Gln Pro Arg Pro Ser His His Gly Val Phe Ala Phe
110                 420                 425                 430
111 Leu Val Thr Pro Ser Pro Tyr Glu Leu Cys Ala Val Pro Arg
112         435                 440                 445
115 <210> SEQ ID NO: 3
116 <211> LENGTH: 10278
117 <212> TYPE: DNA
118 <213> ORGANISM: Human
120 <220> FEATURE:
121 <221> NAME/KEY: misc_feature
122 <222> LOCATION: (1)...(10278)
123 <223> OTHER INFORMATION: n = A,T,C or G
125 <400> SEQUENCE: 3
126 agccttacaa agtgctggga ttacctgcgt gagccaccgg gtccggcctc tttatgtctt 60
127 actgtactgt ctgtcttgaa aagtacttat tatttttgat tggttcatca tttagtctaa 120
128 ttaaaataag agtagtttac acaccacaat tacagtatta taatactctg tttttctgtg 180
129 tgcttactat taccagtgag ttttgtacct ttagatgatt tcttcttgct cattaatatc 240
130 ctttttttt tcagattgaa aaactccctt tagcattct tgtgggatat aggtctggtg 300
131 ttgatgaaat ctcgcagctt ttgtttgtct gggaaggtct ttatttctcc ttcctgttgg 360
132 aaggatattt ttgccagata cgttattcta ggctaaaagt ttttttcct tcagcacttt 420
133 aaatatgtca tgccactccc ccctggcctg taaggtttcc actggaaagg tggctgcccc 480
134 atgtcatgta ttggagctct actgcatgtt atttgtttct tttctcttgc tgcttttagg 540
135 atcctttctt tatccttgac ctttcggagt ttaattatca gatgccttga ggtcgtcttc 600
136 tttgggttaa atctgcttgg tgttctataa acttcttgta caaaaaatca gccaggcatg 660
137 gtggtgggca cctgtaatcc cagctacttg ggaggctgag gcaggagaat cgcttgaacc 720
138 ctggaggtgg aggttgcagt gagccgagat cgcatcattg cactcccacc tgggcgacag 780
139 agcaaaactc cgtctcaaaa aaaaaattat ttgggctcgg tggtgcctgt agtcccagct 840
140 acttgggagg caggaggtcc acttgatgtt gagattgcag tgagccatga tcctgccact 900
141 gcactccggc ccgggcaaca gagtgagacc ctgtctaaag aaaaaataaa aataaaaaag 960
142 caacatatcc taaataaagg atcctccata atgtttccac cagatttcta atcagaaaca 1020
143 tggaggccag gaagcagtgg agaatgacga ccctcaggca gccctggagg atgctgtcac 1080
144 aggctggggc aagggccttc aggctaccaa ctgggagctc tgggaacagc cctgttgcaa 1140
145 acaggaagtc atggcccggc cagagcccag aatgtgggct gagctgggat ccatgtgaca 1200
146 gctttgaggc tcaccgggag cagcctctgg acaggagagg tcccatccag gaaacctcgg 1260
147 gcatggctgg gaagtggggt acttggtgcc gggtctgtat gtgtgtgtga ctggtgtgtg 1320
148 tgagagagaa tgtgtgccct gagtgtcagt gtgagtctgt gtatgtgtga atattgtctt 1380
149 tgtgtgggtg attttctgca tgtgtaatcg tgtccctgca agtgtgaaca agtggacaag 1440
150 tgtctgggag tggacaagag atctgtgcac catcaggtgt gtgcatagcg tctgtgcatg 1500
151 tcaagagtgc aaggtgaagt gaagggacca ggcccatgat gccactcatc atcaggagct 1560
152 ctaaggcccc aggtaagtgc cagtgacaga taagggtgct gaaggtcact ctggagtggg 1620
153 caggtggggg tagggaaagg gcaaggtcat gttctggagg aggggttgtg actacattag 1680
154 ggtgtatgag cctagctggg aggtggatgg ccgggtccac tgagaccctg gttatcccag 1740
155 aagcctgtgt gggcttgggg agcttggagt ggggagaggg ggtgacttct ccgaccaggc 1800
156 ctttctacca ccctaccctg ggtaagggcc tggagcagga agcagcggca aggacctctg 1860
157 gagcagccca tacctgccct ggcctgactc tgccactggc agcacagtca acacagcagg 1920
158 ttcactcaca gcagagggcg aaggccatca tcagctccct ttataaggga agggtcacgc 1980
159 gctcggtgtg ccgagagtgt cctgcctggt cctctgtgcc tggtgggggtg ggggtgccag 2040
```

```
160 gtgtgtccag aggagcccag ttggtagtga ggcagccatg gggctagaag cactggtgcc 2100
161 cctggccatg atagtggcca tcttcctgct cctggtggac ctgatgcacc ggcaccaacg 2160
162 ctgggctgca cgctacccgc caggtcccct gccactgccc gggctgggca accttgctgc 2220
163 atgtggactt ccagaacaca ccatactgct tcgaccaggt gagggaggag gtcctggagg 2280
164 gcggcagagg tcctgaggat gccccaccac cagcaaacat gggtggtggg ttaaaccaca 2340
165 ggctggatca gaagccaggc tgagaagggg aagcaggttt gggggacgtt cctggggaag 2400
166 gacatttata catggcatga aggactggat tttccaaagg ccaaggaaga gtagggcaag 2460
167 ggcctggagg tggagctgga cttggcagtg ggcatgcaag cccattgggc aacatatgtt 2520
168 atggagtaca aagtcccttc tgctgacacc agaaggaaag gccttgggaa tggaagatga 2580
169 gttagtcctg agtgccgttt aaatcacgaa atcgaggatg aaggggtgc agtgacccgg 2640
170 ttcaaacctt ttgcactgtg ggtcctcggg cctcactgct caccggcatg gaccatcatc 2700
171 tgggaatggg atgctaactg gggcctctcg gcaattttgg tgactcttgc aaggtcatac 2760
172 ctgggtgacg catccaaact gagttcctcc atcacagaag gtgtgacccc cacccctgcc 2820
173 ccacgatcag gaggctgggt ctcctccttc cacctgctca ctcctggtag ccccgggggt 2880
174 cgtccaaggt tcaaatagga ctaggacctg tagtctgggg tgatcctggc ttgacaagag 2940
175 gccctgaccc tccctctgca gttgcggcgc cgcttcgggg acgtgttcag cctgcagctg 3000
176 gcctggacgc cggtggtcgt gctcaatggg ctggcggccg tgcgcgaggc gatggtgacc 3060
177 cgcggcgagg acacggccga ccgccccgcct gcgcccatct accaggtcct gggcttcggg 3120
178 ccgcgttccc aaggcaagcg gcggtggggg acagagaccg cgtttccgtg ggccccgggt 3180
179 ggacagtgac cgtagcccaa gcagcgccga caggcgtgg ggtcctggac gtgaaacaga 3240
180 gataaaggcc agcgagtggg ctgaggacag tgggccagga aaccacctgc acggggagg 3300
181 tgcgagtctg tgggctggga ggggcgggg ctactgccca gacccgccag aagcccggtg 3360
182 ggcgagggctg atgcgtcgaa gtggcggtgg cggggaccgc gcctatgctg cgggctcagt 3420
183 gtgggcggga cgggcgggat cttccttgag tggaaaggtg gtcagggtgg gcagagacga 3480
184 ggtggggcca aaccccgccc caggcagggg agcaatgtgg gtgagcaaag agtgggccct 3540
185 gtgcccagct ggaccgggct agggactgcg ggagaccttg tggagcgcca ggggttggagt 3600
186 gggtggcgga gggtggggcc aaggccttca tggcaacgcc cacgtgtccg tcccgcccc 3660
187 aggggtgatc ctgtcgcgct atgggcccgc gtggcgcgag cagaggcgct tctccgtgtc 3720
188 caccttgcgc aacttgggcc tgggcaagaa gtcgctggag cagtgggtga ccgaggaggc 3780
189 cgcctgcctt tgtgccgctt cgccgaccaa gccggtgggt gatgggcaga agggcacaaa 3840
190 gcgggaactg ggaaggcggg ggacggagaa ggcaacccct tacccgcatc tccccacccc 3900
191 caggacgccc ctttcgcccc aacggcctct tggacaaagc cgtgagcaac gtgatcgcct 3960
192 ccctcacctg cgggcgcgc ttcgagtacg acgaccctcg cttcctcagg ctgctggacc 4020
193 tagctcagga gggactgaag gaggagtcgg gctttctgcg cgaggtgcgg agcgagagac 4080
194 cgaggagtct ctgcagggcg agctcctgag aggtgccggg gctggactgg ggcctccgaa 4140
195 gggcaggatt tgcatagatg ggttttgggaa aggacattcc aggagaccccc actgtaagaa 4200
196 gggcctggag gaggagggga catctcagac atggtcgtgg gagaggtgtg cccgggtcag 4260
197 ggggcaccag gagaggccaa ggactctgta cccccgtcca cgttggagat ttcgatttta 4320
198 ggtttctcct ctgggcaagg agagagggtg gaggctggca cttggggagg gacttggtga 4380
199 ggtcagtggt aaggacaggc aggccctggg tctacctgga gatggctggg gcctgagact 4440
200 tgtccaggtg aacgcagagc acaggaggga ttgagacccc gttctgtctg gtgtaggtgc 4500
201 tgaatgctgt ccccgtcctc ctgcacatcc cagcgctggc tggcaaggtc ctacgcttcc 4560
202 aaaaggcttt cctgacccag ctggatgagc tgctaactga gcacaggatg acctgggacc 4620
203 cagcccagcc accccgagac ctgactgagg ccttcctggc aaagaaggag aaggtgagag 4680
204 tggctgccac ggtggggggc aagggtggtg ggttgaacgt cccaggagga atgaggggag 4740
205 gctgggcaaa aggttggacc agtgcatcac ccggcgagcc gcatctgggc tgacaggtgc 4800
206 agaattggag gtcatttggg ggctaccccg ttctatcccc tgagtatcct ctcggccctg 4860
207 ctcaggccaa ggggagccct gagagcagct tcaatgatga gaacctgcgc atagtggtgg 4920
208 gtaacctgtt ccttgccggg atggtgacca cctcgaccac gctggcctgg ggcctcctgc 4980
```

```
209 tcatgatcct acacctggat gtgcagcgtg agcccagctg gggcccaagg cagggactga 5040
210 gggaggaagg gtacagctgg gggcccctgg gcttagctgg gacacccggg gcttccagca 5100
211 caggcgtggc caggctcctg taagcctaac ttcctccaac acaggaggaa ggagagtgtc 5160
212 ccctgggtgc tgacccattg tggggacgca tgtctgtcca gtccgtgtcc aacaggagat 5220
213 cgacgacgtg atagggcagg tgcggcgacc agagatgggt gaccaggctc acatgcccta 5280
214 caccactgcc gtgattcacg aggtgcagcg ctttggggac atcatccccc tgagtgtgac 5340
215 ccatatgaca tcccgtgaca tcgaagtaca gggcttccgc atccctaagg taggcctggc 5400
216 gccctcctca ccccagctca gcaccagcac ctggtgatag cccagcatg gctactgcca 5460
217 ggtgggccca ctctaggaac cctggccacc tagtcctcaa tgccaccaca ctgactgtcc 5520
218 ccacttgggt ggggggtcca gagtataggc agggctggcc tgtccatcca gagcccccgt 5580
219 ctagtgggga gacaaaccag gacctgccag aatgttggag gacccagcgc ctgcagggag 5640
220 aggggggcagt gtgggtgcct ctgagaggtg tgactgcgcc ctgctgtggg gtcggagagg 5700
221 gtactgtgga gcttctcggg cgcaggacta gttgacagag tccagctgtg tgccaggcag 5760
222 tgtgtgtccc ccgtgtgttt ggtggcaggg gtcccagcat cctagagtcc agtccccact 5820
223 ctcaccctgc atctcctgcc cagggaacga cactcatcac caacctgtca tcggtgctga 5880
224 aggatgaggc cgtctgggag aagcccttcc gcttccaccc cgaacacttc ctggatgccc 5940
225 aggccacctt tgtgaagccg gaggccttcc tgccttttctc agcaggtgcc tgtggggagc 6000
226 ccggctccct gtcccccttcc gtgggagtctt gcaggggtat cacccaggag ccaggctcac 6060
227 tgacgcccct cccctcccca caggccgccg tgcatgcctc ggggagcccc tggcccgcat 6120
228 ggagctcttc ctcttcttca cctccctgct gcagcacttc agcttctccg tggccgccgg 6180
229 acagccccgg cccagccact ctcgtgtcgt cagctttctg gtgaccccat ccccctacga 6240
230 gctttgtgct gtgccccgct agaatggggt acctagtccc cagcctgctc cctagccaga 6300
231 ggctctaatg tacaataaag caatgtggta gttccaactt gggtcccctg ctcacgccct 6360
232 cgttgggatc atcctcctca gggcaacccc accctgcct cattcctgct taccccaccg 6420
233 cctggccgca tttgagacgg gtacgttgag gctgagcaga tgtcagttac ccttgcccat 6480
234 aatcccatgt ccccactga cccaactctg actgcccaga ttggtgacaa ggactacatt 6540
235 gtcctggcat gtggggaagg ggccagaatg ggctgactag aggtgtcagt cagccctgga 6600
236 tgtggtggag agggcaggac tcagcctgga ggcccatatt tcaggcctaa ctcagcccac 6660
237 cccacatcag ggacagcagt cctgccagca ccatcacaac agtcacctcc cttcatatat 6720
238 gacaccccaa aatggaagac aaatcatgtc agggagctat atgccagggc tacctcccag 6780
239 ggctcagtcg gcaggtgcca gaacattccc tgggaaggcc ccaggaaaac ccaggaccga 6840
240 gccaccgccc tcagcctgtc accttgtgtc caaaattggt gggttcttgg tctcactgac 6900
241 ttcaagaatg aagccgtgga ccctcacggt gagtgttaca gttcttaaag atggtgtgtt 6960
242 cagagtttgt tccttctgat gttaagacgt gttcagagtt tcttccttct ggtgggtgcg 7020
243 tggtcttgct ggcttcagga gtgaagctgc agaccttcac agtgagtgtt acggctctta 7080
244 aggctgcacg tacggagttg ttcattcttc ctggtgggtt tgtggtctca ctggcctcag 7140
245 gagtgaaact gcagtccttc cagtgttaca actcataaag gcagtgtgga cccaatgagg 7200
246 gagcagcagc agcaagactt actgcaaaca gcaaagaat gatggcaacc aggttgccgc 7260
247 tgctacttca ggcagcctgc ttttattccc ttatctgacc cccacccaca tcctgctgat 7320
248 tggcccattt tacagacagt ggattggtcc acttacagag agctgattgg tgcatttaca 7380
249 atccctgagc tagacacaga gtactgattg gtatatttac aaaccttgag ctagacacag 7440
250 agtgctgaat ggtgtatta caatcccta gctagacata aaggttgtcc cagtccccac 7500
251 tagattagct agatagagta gacagagagc actgattggt gcgtttacaa accttgagtt 7560
252 agacacaggg tgctgactgg tgtgtttaca aaccttgagc tagacacaga gtgctgattg 7620
253 gtgtatttac aatctttttag ctagaaataa aggttcccca agtccccacc agattagcta 7680
254 gatagagtgc taattggtgc atgcacgaac ccggagctag acacagagtg ctgattggtg 7740
255 catatacaat cctctggcta gacataaaag ttctccaagt ccccacctga ctcaggagcc 7800
256 cagccagctt cgcctagtgg atcctatgcc agggccacag gcagagctgc ctgctagtcc 7860
257 cacaccgggc acctgtactc ctcagccctt gggcagtgga cgggaccagg tgccgtggag 7920
```

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/820,788

DATE: 04/10/2001
TIME: 15:48:02

Input Set : A:\Seqlist.txt
Output Set: N:\CRF3\04102001\I820788.raw

L:14 M:270 C: Current Application Number differs, Replaced Current Application No
L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:279 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:280 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:281 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3